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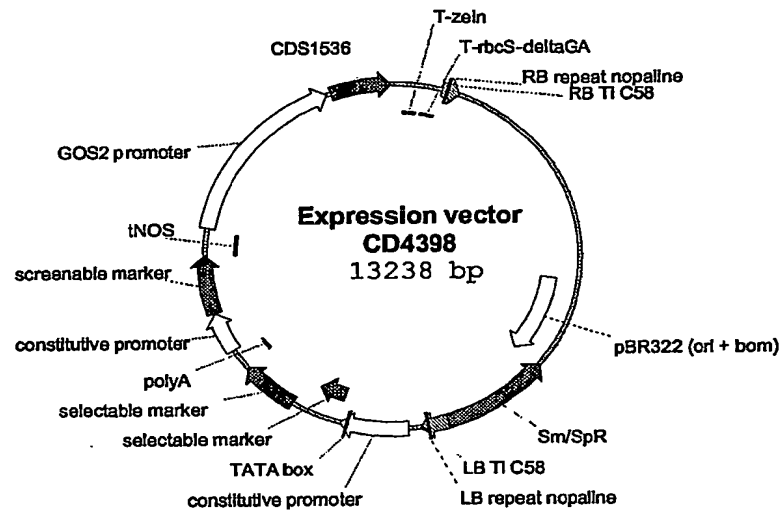


FIGURE 1

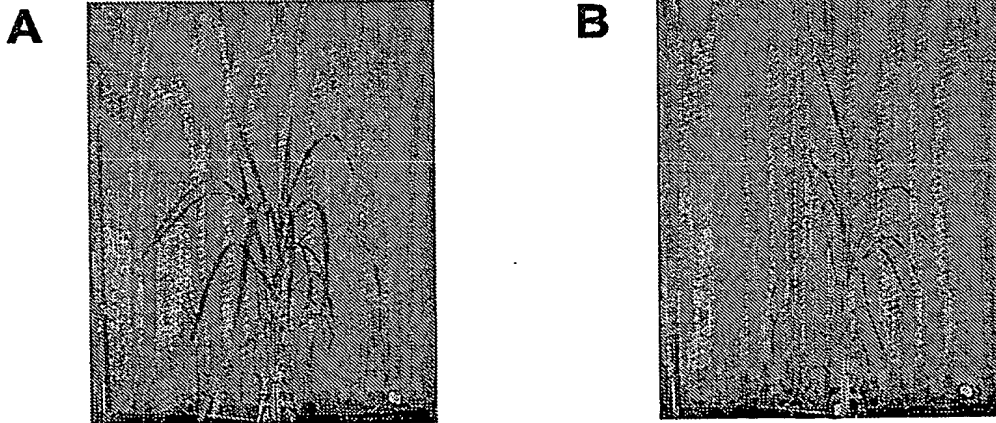


FIGURE 2

### FIGURE 3

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## ORTHOLOGS OF STZ in OTHER PLANT SPECIES

SEQ ID NO 10: Dg\_AF119050\_1 *Datisca glomerata* zinc-finger protein 1 (zfp1) mRNA, complete cds

GGCACGAGGACAAATTCTCTCTATCCTCTGAATATCTTTGGTTTGTGAACTGAGAAGCTA  
TTAGATGGCTCTAGAAGCGCTCAACTCTCCGACCACAGCTACGCCGGTGTTCCTACTACGACG  
ACCCAGCTTGAATTACCTTGAGCCATGGACCAAGCGTAAGCGTTCCAAGCGTACGCGCTTA  
GATAGCCCCATACCGAGGAAGGTACCTTGCTTTCTGCCTCATCATGCTCGCTCGTGGCCGC  
GTTGCCTCTGCAAATCGACGGGATTCTCAGTCTTCCATTTCAGATTTCAGCCTGAAGCAACGAC  
TTCCGGCTACCAAAGTCAGTTATAAGTGCTCTGTGTGCGATAAGGCCCTTTTCGTCTTATCAGG  
CTTTGGGTGGGCACAAGGCCAGCCACAGAAAGCTCGCTGGCGGCGAAGATCAATCGACTTCC  
TTTGCCACCACGAATTCAGCCACCGTCACTACCACCACAGCCTCCGGAGGTGGTGGCAGGTC  
TCATGAGTGTTCTATTTGCCACAAATCGTTCCCGACTGGCCAGGCCCTTGGGTGGTCACAAGC  
GCTGCCACTACGAAGGCAGTATCGGCGGCAATAGTATTCACCACCACAACAATACCACCAAC  
AGCGGAAGCAACGGTGGCATGAGCATGACCTCCGAAGTAGGTTCCACACACACAGTCAGCCA  
CAGTCACCGTGACTTCGATCTCAACATCCCGGCCTTGCCGGAGTTTCGGTCAATTTCTTCA  
TATCCGGGGATGACGAGGTCGAGAGTCTCATCCGGCCAAGAAACCCCGTATATTGATGAAA  
TAAAACATTTCTCAAGATCACTGAACAGGCTTTAGTTTCTTTATAGGAGGAGATTTAAAAA  
AGTAGTATCTCTCTTTCTTTATCCGTAGGATAATTAATATATTTTCGTGTACATAAATTTGTA  
GTTCTTTAACACACTCTGTTTCATTTTCTTGCTTTGCTCAACTTTGTATTGGTTATTTTCATT  
ATGAAAATTCAATT

SEQ ID NO 11: Dg\_AF119050\_1 *Datisca glomerata*, STZ ortholog, protein

MALEALNSPTTATPVFHYDDPSLNYLEPWTKRKRSKRTRLDSPHTEEEYLAFCLIMLARGRV  
ASANRRDSQSSIQIQPEATTSATKVSYSKSCVCDKAFSSYQALGGHKASHRKLGGEDQSTSF  
ATTNSATVTTTTASGGGGRSHECSICHKSFTPGQALGGHKRCHYEYSIGGNSIHNNNTNS  
GSNGGMSMTSEVGSTHTVSHSHRDFDLNIPALPEFRSNFFISGDDEVESSHPAKKPRILMK

SEQ ID NO 12: Gm\_T09602\_U68763.1\_GMU68763 *Glycine max* (soybean) probable zinc finger protein SCOF-1 mRNA, complete cds

AAAATTCTCACTCTCTCTCATCTCGAGATCATAGTATCATATTCAATATCATTTTCATACC  
AAACACATGGCTTTGGAAGCTCTCAACTCACCAACAACAACCGCTCCATCTTTTCCCTTTGA  
CGACCCAACTATTCCATGGGCGAAACGAAAACGTTCAAAGCGTTCTCGCGACCATCCTTCTG  
AAGAAGAGTACCTCGCCCTCTGCCTCATCATGCTCGCTCGCGGCGGCACCACCACCGTCAAC  
AACCGCCACGTCAGCCCTCCGCCGCTACAGCCACAGCCACAGCCGACACCAGATCCTTCCAC  
CAAGCTCAGTTACAAATGCTCCGTTTGCGACAAGAGCTTCCCTCTTACCAAGCGCTCGGTG  
GACACAAGGCCAGTCACCGGAAACTCGCCGCGCGCCGCGGAAGACCAACCCCCCAGCACCACC  
ACTTCCTCCGCCGCGGCCACCAGCTCCGCCCTCCGGAGGTAAGGCCCATGAGTGCTCCATTTG  
CCACAAATCCTTCCCCACCGGACAGGCCCTTGCGGGACACAAACGTTGTCACTACGAAGGTA  
ACGGTAACGGAAATAACAACAACAGTAACAGCGTTGTACCGTCGCCTCGGAAGGCGTGGGC  
TCCACCCACACTGTCAGTCACGGCCACCACCGCGACTTCGATCTCAACATCCCGGCCCTTTCC  
GGATTTTTTCGACCAAGGTCGGAGAAGACGAGGTTGAGAGCCCTCACCTGTCAATGAAGAAGC  
CTCGCCTCTTCGTCACTATATATTTTGGAGAATTTTGGAGAGTTACTATAATTTGATTTGTAC  
ATAGTACTTGGAAGTTTTGTTGGACCGTACCGGACCCAGTCTCTGGTTGAGGTTGTACTTT  
CACAAACAGTGGCAGATTTGCAATTCAATTCAATTTATTTGTTTATTTTAAAAAAAAAAAAAA  
AAAA

FIGURE 3 (continued)

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**SEQ ID NO 13:** Gm\_T09602 Glycine max (soybean) probable zinc finger protein SCOF-1, STZ otholog, protein  
 MALEALNSPTTTAPSFPPFDDPTIPWAKRKRKRSRDHPSEEEYLALCLIMLARGGTTTVNNR  
 HVSPPPLQPQPQPTPDPSTKLSYKCSVCDKSFPSYQALGGHKASHRKLGAEDQPPSTTTS  
 SAAATSSASGGKAHECSICHKSFPTGQALGGHKRCHYEGNGNGNNNNNSNSVTVVASEGVGST  
 HTVSHGHHRDFDLNIPAFPDFSTKVGEDEVESHPVMKKPRLFVIPKIEIPQFO

**SEQ ID NO 14:** Ms\_CAB77055\_Y18788.1\_MSY18788 Medicago sativa putative TFIIIA (or kruppel)-like zinc finger protein mRNA  
 AATTCGGCAGAGAAATAACCACTTCTCTCTCAAAACCTCCTTTTGCCTTTTGCTTCTACTT  
 TCACTTGCCTAACGCTAACTAACTCTTCTCGAGTGTTCTTCTTTTCATCATATGGCTATGGA  
 AGCACTTAACTACCCACCACTGCTACTCCTTTTACACCCCTTTGAGGAACCAATCTGAGTT  
 ATCTTGAAACACCGTGGACGAAAGGTAAACGATCAAAGCGTTCTCGCATGGATCAATCTTCA  
 TGCCTGAAGAAGAGTATCTCGCTCTTTGTCTCATCATGCTTGCTCGCAGCGGTAACAACAA  
 CGACAAAAGTCTGATTCGGTGGCGACGCCGCTAACCAACCGTTAAACTCAGTCACAAATGCT  
 CAGTCTGCAACAAAGCTTTCTCATCTTATCAAGCCCTAGGTGGACACAAAGCCAGTCACCGG  
 AAAGCTGTTATGTCCGCAACCACCGCTGAAGATCAGATCACCACCACTTCATCCGCCGTGAC  
 TACCAGCTCTGCTTCCCAACGGTAAGAACAAGACTCATGAGTGTTCCATCTGTACAAATCCT  
 TCCCTACTGGACAGGCTTTGGGAGGACACAAGCGTTGTCACTACGAAGGCAGCGTTGGTGCC  
 GGTGCCGGTGCTGGAAGTAACGCTGTAAGTGCCTCTGAAGGAGTTGGATTGTACACAGCCA  
 CCACCGTGATTTTGATCTTAACCTCCCGGCTTTTCCGGACTTTTCAAAGAAGTTTTTCGTGG  
 ATGACGAGGTTTTTAGTCCTTTACCTGCTGCAAAGAAGCCCTGTCTTTTCAAGCTGGAAATT  
 CCTTCTCATTACTGATCAATAATAGATCCAATTTTATTGTTATTATTATTAATAATTATTAT  
 CGCTTAGGGCATAAGTTATTTTCTTTTTCTTTCAATTATTTTCGGATCAATTTGTTCTGTACA  
 TACAAATTGGGATTGGTTTTAGAATTTAGGACGGTTGTAGACAATGGAAATTCAATTCAATT  
 ATTTAATTTTGTGT

**SEQ ID NO 15:** Ms\_CAB77055\_Medicago sativa putative TFIIIA (or kruppel)-like zinc finger protein, STZ otholog, protein  
 MAMEALNSPTTATPFTPFEENLSYLETPWTKGKRKRSRMDQSSCTEEYALCLIMLARS  
 GNNNDKKS DSVATPLTTVKLSHKCSVCNKAFFSSYQALGGHKASHRKAVMSATTAEDQITTS  
 SAVTTSSASNGKNKTHECSICHKSFPTGQALGGHKRCHYEGSVGAGAGAGSNAVTASEGVGL  
 SHSHHRDFDLNLPAFPDFSKKFFVDDEVFSPLPAKKPCLEFKLEIPSHY

**SEQ ID NO 16:** Nt\_AAC06243\_AF053077 Nicotiana tabacum osmotic stress-induced zinc-finger protein (zfp) mRNA, complete cds  
 TTTTCCCTCGAATTTGATAACTAAAGAGAATATTATGACTCTTGAAGCTTTGAAGTCACCTA  
 CGGCGGCAACGCCGACTCTACCACCACGCTATGAAGATGATGATGAAATTCATAATTTGGAT  
 TCTTGGGCTAAAGGAAAACGATCAAAACGGCCCCGATTGATGCCCCACCGACTGAAGAAGA  
 GTATTTAGCCCTCTGTCTCATCATGCTCGCTCGCAGCGGAACCGGAACCAAGACCGGTTTAA  
 CTGATGCTACTACTTCCCAACAACCTGCCGATAAAAAAACCGCCGAGTTGCCGCCGGTTTCA  
 AAGAAAGAGGTGGCAACAGAGCAAGCAGAGCAATCTTACAAGTGTAGCGTGTGTGACAAGGC  
 TTTTCTTCTTATCAAGCACTCGGTGGGCATAAAGCAAGTCACCGTAAACTACTACTACTG  
 CTACCGCCGCTCTGATGATAACAATCCTTCAACTTCAACTTCCACTGGCGCCGTTAATATC  
 TCTGCTCTTAATCCAACCTGGTTCGTTTACACGCTGTCTTATTTGCCACAAGGCTTTTCTTAC  
 TGGCCAAGCTTTGGGTGGGCACAAGCGCCGCACTATGAAGGCAAACTCGGTGGTAACAGCC  
 GCGACTTAGGCGGCGGCGGCGGCGGCGGTCATAGTGGAAGCGTCTTGACTACTTCAGACGGC

FIGURE 3 (continued)

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GGCGCGTCGACTCACACGCTACGTGACCTTGACCTGAACATGCCTGCTTCGCCGGAATTGCA  
 ACTGGGTCTGAGTATTGATTGTGGACGGAAAAGTCAACTGTTGCCGATGGTCCAAGAGGTGG  
 AAAGTCCTATGCCTGCAAAGAAACCGCGTTTATTGTTTTCGTTGGGTGAAACTTCTTTAGG  
 GGAATTGAATTGATTGTGTTTACGCCAAATTAGTAAATTGGTTCATGTGATTTTATTTTATG  
 GAAAAGGAATTATTGATTGTTTTACCCGTTTATTCTTAGGGTGGTATTATGTACAGGGAGTG  
 AATCATTCAATTGGTTTTACACTTTCTTAATTATATATTCTTTTTTTTTTACACATAAAAAAAA  
 AAAAAA

**SEQ ID NO 17:** Nt\_AAC06243\_Nicotiana tabacum osmotic stress-  
 induced zinc-finger protein, STZ ortholog, protein  
 MTLEALKSPTAATPTLPPRYEDDDEIHNLD SWAKGRSKRPRIDAPPTEEYLALCLIMLAR  
 SGTGTRTGLTDATTSQQPADKKTAE LPPVHKKEVATEQAEQSYKCSVCDKAFSSYQALGGHK  
 ASHRKTTTTATAASDDNNPSTSTSTGAVNISALNPTGRSHVCSICHKAFPTGQALGGHKRRH  
 YEGKLGGNSRDLGGGGGGHSGSVLTTS DGGASTHTLRDFDLNMPASPELQLGLSIDCGRKS  
 QLLPMVQEVESPMPPAKKPRLLFSLG

**SEQ ID NO 18:** Os\_AF332876 Oryza sativa zinc finger  
 transcription factor ZF1 mRNA, complete cds  
 AATTCGGCAGCAGGCCACACAGCAACCAGCCAGCTGCCACACTAGCTTGAGGCGAGCGAGCG  
 AAGCTTAGCTAGCGGATAGAACAAGTCGTCGATTGCTTGCTTTTGTGAATTGCGGTGG  
 AAGCATGTCGAGCGCGTCGTCATGGAAGCGCTCCACGCCGCGGTGCTCAAGGAGGAGCAGC  
 AGCAGCACAGGTGGAGGAGGCGACGGTCGTGACGAGCAGCAGCGCCACGAGCGGGGAGGAG  
 GGCGGACACCTGCCCGAGGGGTGGGCGAAGCGGAAGCGGTGCGCGCCGCGCAGCGATCGGAGGA  
 GGAGAACCTCGCGCTCTGCCTCCTCATGCTCGCCCGCGCGGCCACCACCGCTCCAGGCGC  
 CGCCTCCGCTCTCGGCTTCGGCGCCCCCGCCGGCAGGTGCGGAGTTCAAGTGCTCCGCTCTGC  
 GGCAAGTCCTTACGCTCCTACCAGGCGCTCGGCGGCCACAAGACGAGCCACCGGGTCAAGCT  
 GCCGACTCCGCGCGCAGCTCCCGTCTTGCTCCCGCCCCCGTCCGCGCCTTGCTGCCTTCCG  
 CCGAGGACCGCGAGCCAGCCACGTCATCCACCGCCGCGTCTCCGACGGCATGACCAACAGA  
 GTCCACAGGTGTTCCATCTGCCAGAAGGAGTCCCCACCGGGCAGGCGCTCGGCGGGCACAA  
 GAGGAAGCACTACGACGGTGGCGTAGGCGCCGCGCGCGCATCTTCAACCGAGCTCCTGG  
 CCACGGTGGCCGCCGAGTCCGAGGTGGGAAGCTCCGGCAACGGCCAGTCCGCCACCCGGGCG  
 TTCGACCTCAACCTCCCGGCCGTGCCGGAGTTCGTGTGGCGGCCGTGCTCCAAGGGCAAGAA  
 GATGTGGGACGAGGAGGAGGAGTCCAGAGCCCCCTCGCCTTCAAGAAGCCCCGGCTTCTCA  
 CCGCGTAATTCAGCAGCTGCACGGATCCGATCCGTCAGAGTTTTTGTCTAGGGAGTGAAATT  
 CAGTCGAAACACACTATTTCGTTGATTTCGTTTTGTGCCGCTATTGTTTAATTGTTCCGTGCTT  
 TTGTACAGAGCAAGCGAGTGATACATAGCCATACATACAGTCATACAGATATAGGTCTAGCT  
 CTTCTTGGTTCTTTGTAACTGGAAGTGTACCTGTATCTTTTACACTTTGTCTTTGACA  
 GTCATATATTGTAGACCAAAAAAAAAAAAAAAAAA

**SEQ ID NO 19:** Os\_AF332876 Oryza sativa seedling zinc finger  
 transcription factor ZF1, STZ ortholog, protein  
 MSSASSMEALHAAVLKEEQQHEVEEATVVTSSSATS GEEGGHLPQGWAKRKRSRRQRSEE  
 NLALCLIMLARGGHRVQAPPLSASAPPPAGAEFKCSVCGKSFSSYQALGGHKTSRHKLP  
 TPPAAPVLAPAPVAALLPSAEDREPATSSSTAASSDGMTNRVHRC SICQKEFPTGQALGGHKR  
 KHYDGGVGAGAGASSTELLATVAAESEVGSSNGQSATRAFDLNLPAVPEFVWRPCSKGKKM  
 WDEEEEVQSPLAFKKPRLLTA

FIGURE 3 (continued)

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**SEQ ID NO 20:** Ph\_BAA05079\_D26086.1 [Petunia x hybrida].  
PETZFP4 zinc-finger protein gene  
TTCACCTCACCAAAACAACCTTCTCTACCTCTTCTACTTGACATTCAAATTCTTTTCACTACTA  
CTTATCTCTACTAATCTTGATTTCGATTTTAGTAAATCAAACAAGAGAATCTTTTCAGTAATA  
CAAACAAGAAAATTTCTCTCTATACTTGATTGAGTTTAGTAAGGCAAACAAGAAAACATC  
**ATGGCACTTGAAGCATTGAATTCTCCAAC**TACAACAACACCACCATCATTCCAATTGAGAA  
CAACGGGCTTAAGTACCTTGAGAGTTGGACAAAAGGTAAAAGATCAAAAAGGCAACGCAGCA  
TGGAACGACAGTGTACTGAAGAAGAGTATTTAGCACTTTGTCTTATCATGCTAGCACGTAGC  
GATGGTTCTGTAAATAACTCACGGTCTCTACCACCACCACCCTACCACCATCAGTTCCAGT  
AACGTCGCAAATAAACGCGACGTTATTGGAACAGAAGAATTTGTACAAGTGTTCCGTTTGTG  
GTAAAGGGTTTGGGTCTTATCAAGCTTTAGGTGGACATAAAGCAAGTCACCGGAAACTTGTG  
AGCATGGGAGGAGATGAACAATCTACTACTTCCACTACTACTAAACGTAACGGGAACTAGTTC  
CGCTAACGTTAACGGTAACGGAAGAACTCACGAATGTTCAATTTGTCACAAGTGCTTTCCTA  
CTGGACAAGCTTTAGGTGGTCATAAAGGTGCCACTATGACGGTGTTAACGGTAACGGTAAC  
GGAAGTGTAAGTGTTGGGGTGACGTCACTGAAGGTGTGGGGTCCACTATTAGTCATCCAGG  
TGACTTTGACTTGAATATTTCCCGGTTGCCGGAGTTTGGCCGGGATTTGGTTCCGGCGAGG  
ATGAGGTGGAGAGTCTCATCCAGCAAAGAAGTCAAGGCTATCTCTTCCACCTAAACTTGAA  
TTATTCAAAGGATTAT**AGAGGGAATATTGATT**TGTTACAGGAAGATTTATTAGGATTCACGA  
ATTTTTTGTGACTAGTTTATGTAATAT

**SEQ ID NO 21:** Ph\_BAA05079 [Petunia x hybrida] zinc-finger  
protein, STZ ortholog, protein  
MALEALNSPTTTTPPSFQFENNGLKYLESWTKGKRKRQRSMEROCTEEYLAALCLIMLARS  
DGSVNNRSRLPPPLPPSPVPTSQINATLLEQKNLYKCSVCGKGFGSYQALGGHKASHRKL  
SMGGDEQSTTSTTTNVTGTSSANVNGNGRTHCESICHKCFPTGQALGGHKRCHYDGGNGNGN  
GSVSVGVTSSSEVGSTISHHRDFDLNIPALPEFWPGFSGEDEVESPHPAKKSRLSLPPKLE  
LFKGL

**SEQ ID NO 22:** Ta\_BAA03901 Triticum aestivum gene for zinc-  
finger protein WZF1, complete cds  
**ATGTCGTCGTCGGCCATGGAAGCGCTCCACGCCCTGATCCCGGAGCAGCACCAGCTGGACGT**  
TGAGGCGGCTGCGGCTGTCAGCAGCGCCACCAGCGGCGAGGAGAGCGGCCACGTGCTGCAGG  
GGTGGGCCAAGAGGAAGCGATCGCGCCGCCAGCGCTCCGAGGAGGAGAACCCTCGCGCTCTGC  
CTCCTCATGCTCTCGCGCGGCGGCAAGCAGCGTGTTCAGGCGCCGAGCCGGAGTCGTTTCGC  
TGCGCCGGTGCTGCCGAGTTCAAGTGCTCCGTCTGCGGCAAGTCCTTCAGCTCCTACCAGG  
CGCTCGGAGGCCACAAGACGAGCCACCAGGTGAAGCAGCCGTCTCCTCCCTCTGATGCCGCT  
GCTGCCCCACTCGTGGCCCTCCCGGCCGTGCGCGCCATCTGCCGTCCGCCGAGCCGGCCAC  
GTCGTCCACCGCCGCGTCTCCGACGGCGCGACCAACAGAGTCCACAGGTGCTCCATCTGCC  
AAAAGGAGTTCCCGACTGGGCAGGCGCTCGGCGGGCACAAGAGGAAGCACTACGACGGAGGC  
GTGGGCGCCGCCGCTCGTCGACCGAGCTTCTGGCCGCCGCGGCCGCGAGTCTGAGGTGGG  
GAGCACC GGCAACGGGAGCTCCGCCGCCCGGCCCTTCGACCTGAACATTCCGGCCGTGCCGG  
AGTTCGTGTGGAGGCCGTGCGCCAAGGGCAAGATGATGTGGGAGGACGATGAGGAGGTGCAG  
AGCCCCCTCGCCTTCAAGAAGCCTCGGCTTCTCACCGCT**TGA**

FIGURE 3 (continued)

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**SEQ ID NO 23:** Ta\_BAA03901\_WZF1 *Triticum aestivum*, STZ otholog, protein  
 MSSSAMEALHALIPEQHQLDVEAAA AVSSATSGEESGHVLQGWAKRKRSRRQRSEENIALC  
 LLMLSRGGKQRVQAPQPEFSAAPVPAEFKCSVCGKSFSSYQALGGHKTSHRVKQPSPPSDAA  
 AAPLVALPAVAAILPSAEPATSSSTAASSDGATNRVHRCSICQKEFPTGQALGGHKRKHYDGG  
 VGAAASSTELLAAAAAESEVGSTGNGSSAARAFDLNIPAVPEFVWRPCAKGKMMWEDDEEVQ  
 SPLAFKKPRLTLTA

**SEQ ID NO 24:** Ca AF539746 *Capsicum annum* zinc finger protein mRNA, complete cds  
 AAAATCTTCGCTACTTACTTACATCTTCTAGAATAGTCACTAGAACCACTAACTTTATACAA  
 CGGATATCGATATGGCACCTTGAAGCTTTGAATTCCTCAACTGGTACACCAACTCCGCCACCG  
 TTTCAATTTGAGAGCGACGGCCAACAGCTTCGATATATCGAAAACCTGGAGGAAGGGAAAGAG  
 ATCTAAAAGGTCACGCAGCATGGAGCACCAGCCTACTGAGGAAGAATACTTAGCGCTTTGTT  
 TGATCATGCTTGACGTAGCGGTGGCTCCGTTAATCATCAACGATCTCTACCACCGCCGGCT  
 CCGGTGATGAAACTGCACGCGCCGTCGTATCATCGGCGGCGGAGGAGGAGAAGGAGAAGAT  
 GGTGTATAAGTGTTTCGGTTTGTGGTAAGGGATTTGGGTCTTATCAAGCTTTAGGTGGACACA  
 AAGCTAGTCACCGGAAACTCGTACCCGGCGGAGATGATCAGTCAACTACCTCCACAACCACT  
 AACGCAACCGGAACAACAACCTCCGTTAACGGCAACGGCAACAGAAGTGGAAGGACTCACGA  
 GTGTTTCGATTTGTCAAGTGTTTCCCACTGGACAAGCTTTAGGTGGACACAAAAGGTGTC  
 ACTACGACGGCGGTATCGGTAACGGAAACGCTAACAGTGGCGTTAGTGCTAGCGTTGGAGTG  
 ACGTCATCGGAGGGTGTGGGGTCCACAGTCAGTCACCGGGATTTGACTTGAACATTCCGGC  
 GTTGCCGGAATTCTGGCTGGGATTTGGTTCCGGCGAAGATGAGGTGGAGAGTCCACATCCGG  
 CGAAGAAATCGCGGTTATGTTTGCCTCCAAAATATGAATTATTTCAACATTAATGGGAATTT  
 GATTGTTAGGATTTACTATTTTGGTAGACAAAATTATACTATGTAAGTTTTAATTTTCATTG  
 TGGGTGGGAGCAAAATTTTTAATTTTTTGTCTATAGACCTAGCTAGTTACTAATAGCAAAAA  
 TTCAATTGATTGATTTAAAAA

**SEQ ID NO 25:** Ca AF539746-capsicum annum, STZ otholog, protein  
 MALEALNSPTGTPTPPPFQFESDGOQLRYIENWRKGRSKRSRSMEHQPTEEEEYLALCLIML  
 ARSGSVNHQSRSLPPPAPVMKLHAPSSSSAAEEEEKEKMYKCSVCGKGFGSYQALGGHKASH  
 RKLVPGGDDQSTSTTTNATGTTTSVNGNGNRSGRTHCESICHKCFPTGQALGGHKRCHYDG  
 GIGNGNANSVGSASVGVTSSEGVGSTVSHRDFDLNIPALPEFWLGFSGEDEVESPHPAKKS  
 RLCLPPKYELEFQH

#### PARALOGS OF STZ IN ARABIDOPSIS THALIANA

**SEQ ID NO 26:** gi\_18402298\_ref\_NM\_112848.1 mRNA  
 ACTTCACTCTCTAATTTCTCTCTCTATCTCTCACCATATTCGCGATTAAAACTCTCAAC  
 TTTTCTCTCAAATTTCTGATCCTTTGATCCAACAGTTAGAAGAAGATTCATCTGATCATGGC  
 CCTCGAAGCGATGAACACTCCAACCTCTTCTTTACCAGAATCGAAACGAAAGAAGATTTGA  
 TGAACGACGCCGTTTTTCATTGAGCCGTGGCTTAAACGCAACGCTCCAAACGTCAGCGTTCT  
 CACAGCCCTTCTTCGTCTTCTCCTCACCAGCTCGATCTCGACCCAAATCCCAAGATCAAGA  
 TCTTACGGAAGAAGAGTATCTCGCTCTTTGTCTCCTCATGCTCGCTAAAGATCAACCGTCGC  
 AAACGCGATTTTCATCAACAGTCGCAATCGTTAACGCCGCCGCGCCAGAATCAAAGAACCTTCG  
 TACAAGTGTAACGTCTGTGAAAAGCGTTTCTCTCCTATCAGGCTTTAGGCGGTCAAAAGC  
 AAGTCACCGAATCAAACCAACCGTAATCTCAACAACCGCCGATGATTCAACAGCTCCGA

**FIGURE 3 (continued)**

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CCATCTCCATCGTCGCCGGAGAAAAACATCCGATTGCTGCCTCCGGAAGATCCACGAGTGT  
TCAATCTGTCATAAAGTGTTCGACGGGTCAAGCTTTAGGCGGTCACAAACGTTGTCTACTA  
CGAAGGCAACCTCGGCGGCGGAGGAGGAGGAGGAAGCAAATCAATCAGTCACAGTGGAAGCG  
TGTCGAGCACGGTATCGGAAGAAAGGAGCCACCGTGGATTTCATCGATCTAAACCTACCGGCG  
TTACCTGAACTCAGCCTTCATCACAATCCAATCGTCGACGAAGAGATCTTGAGTCCGTTGAC  
CGGTAAAAAACCGCTTTTGTGACCGATCAGACCAAGTCATCAAGAAAGAAGATTTATCTT  
TAAAAATCTAATACTCGACTATTAATTCTTGTGTGATTTTTTTCGTTACAACCATAGTTTCA  
TTTTCATTTTTTTAGTTACAAATTTTAATTGTTCTGATTTGGATTGAATATTGGTATATTG  
TTAGGGGTTGATAC

**SEQ ID NO 27:** Translation of gi\_18402298\_ref\_NM\_112848.1  
MALEAMNTPPTSSFTRIETKEDLMNDAVFIEPWLKRKRKRQRSHSPSSSSSSPPRSRPSQ  
QDLTEEEYLALCLLMLAKDQPSQTRFHQSQSLTPPPESKNLPYKCNVCEKAFPSYQALGGH  
KASHRIKPPTVISTTADDSTAPTISIVAGEKHPIAASGKIHECSIHKVFPTGQALGGHKRC  
HYEGNLGGGGGGGSKSISHSGSVSSTVSEERSHRGFIDLNLPALPELSLHHNPIVIDEEILSP  
LTGKKPLLLTDHDOVIKKEDLSLKI

**SEQ ID NO 28:** gi\_30680473\_ref\_NM\_120516.3\_mRNA  
AAATCAAATCTTTTCATTTACAATTATCTTTCTCTCAATTTAGAACTTAGTAGCTAGTCTT  
CAAGATAATGGCACTTGAACTCTTACTTCTCCAAGATTATCTTCTCCGATGCCGACTCTGT  
TTCAAGATTCAGCACTAGGGTTTCATGGAAGCAAAGGCAAACGATCTAAGCGATCAAGATCT  
GAATTCGACCGTCAGAGTCTCACGGAGGATGAATATATCGCTTTATGTCTCATGCTTCTTGC  
TCGCGACGGAGATAGAAACCGTGACCTTGACCTGCCTTCTTCTTCGTCTTCACCTCCTCTGC  
TTCTCCTCTTCTCTACTCCGATCTACAAGTGTAGCGTCTGTGACAAGGCGTTTTTCGTCTTAC  
CAGGCTCTTGGTGGACACAAGGCAAGTCACCGGAAAAGCTTTTCGCTTACTCAATCTGCCGG  
AGGAGATGAGCTGTGACATCGTCGGCGATAACCACGTCTGGTATATCCGGTGGCGGGGGAG  
GAAGTGTGAAGTCGCACGTTTGCTCTATCTGTCATAAATCGTTCGCCACCGGTCAAGCTCTC  
GGCGGCCACAAACGGTGCCACTACGAAGGAAAGAACGGAGGCGGTGTGAGTAGTAGCGTGTG  
GAATTCTGAAGATGTGGGGTCTACAAGCCACGTCAGCAGTGGCCACCGTGGGTTTGACCTCA  
ACATACCGCCGATACCGGAATTCTCGATGGTCAACGGAGACGAAGAGGTGATGAGTCTTATG  
CCGGCGAAGAAACTCCGGTTTGACTTCCCGGAGAAACCCTAAACATAAACCTAGGAAAAACT  
TTACAGAATTCATTTTATAGGAAATTGTTTTACTGTATATACAAATATCGATTTTGATTGAT  
GTTCTTCTTCACTGAAAAATTATGATCTTTGTTGTATAATTGATGTTTCTGAAAAGATAT  
AACTTTTTATTGTTTCACACGTATCAAAATTTGCTTGGATACATCA

**SEQ ID NO 29:** Translation of gi\_30680473\_ref\_NM\_120516.3  
MALETLTSPRLSSPMPTLFQDSALGFHSGKGRSKRSRSEFDRQSLTEDEYIALCLMLLARD  
GDRNRDLPLPSSSSPPLPLPTPIYKCSVCDKAFSSYQALGGHKASHRKSESLTQSAGGD  
ELSTSSAITTSIGISGGGGSVKSHVCSICHKSFATGQALGGHKRCHYEGKNGGVSSSVSNS  
EDVGSTSHVSSGHRGFDLNIPIPEFSMVNGDEEVMSMPAKKLRFDFPEKP

FIGURE 3 (continued)



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**SEQ ID NO 30:** gi\_30693252\_ref\_NM\_114853.2\_mRNA

ATGGCTCTCGACACTCTCAATTCTCCACCTCCACCACCACAACCACCGCTCCTCCTCCTTT  
CCTCCGTTGCTTCGACGAAACCGAGCCCGAAAACCTCGAATCATGGACCAAAAGAAAACGTA  
CAAAACGTCACCGTATAGATCAACCAACCCCTCCTCCTTCTGAAGAAGAGTATCTCGCTCTT  
TGCCTCCTTATGCTCGCTCGTGGCTCCTCCGATCATCACTCTCCACCGTCGGATCATCACTC  
TCTTTCTCCACTGTCCGATCATCAGAAAGATTACAAGTGTTCGGTCTGTGGCAAATCTTTCC  
CGTCTTACCAAGCGTTAGGTGGACACAAAACAAGTCACCGGAAACCGGTTAGTGTGATGTT  
AATAATAGTAACGGAACCGTTACTAATAACGGAAATATTAGTAACGGTTTAGTTGGTCAAAG  
TGGGAAGACTCATAACTGCTCTATATGTTTTAAGTCGTTTCCCTCTGGTCAAGCATTGGGTG  
GTCACAAACGTTGTCACTATGATGGTGGTAACGGTAACAGTAACGGTGACAATAGCCACAAG  
TTTGACCTAAATTTACCGGCTGATCAAGTTAGTGATGAGACAATTGGAAAAAGTCAACTCTC  
CGGTGAAGAAACAAAGTCGGTGTGTGATTATTATTTTTTACCGATCGGGATTAGCTAG  
TGGTTGATCATTAGCTGAGTCTGTAATGAAAATGAT

**SEQ ID NO 31:** Translation of gi\_30693252\_ref\_NM\_114853.2

MALDTLNSPTSTTTTTAPPFLRCLDETEPENLESWTKRKRTKRHRIDQNPSPSEEEYLAL  
CLIMLARGSSDHHSPPSDHHSLSPLSDHQDKDYKCSVCCKSFPSYQALGGHKTSHRKPVSVDV  
NNSNGTVTNNGNISNGLVGQSGKTHNCISICFKSFPSGQALGGHKRCHYDGGNGNSNGDNSHK  
FDLNLPAQVSDETIGKSQLSGEETKSVL

**SEQ ID NO 32:** gi\_30694224\_ref\_NM\_123683.2\_mRNA

AAATTTTCTATAGCAATGGCGCTTGAAGCTCTTAATTCACCAAGATTGGTCGAGGATCCCTT  
AAGATTCAATGGCGTTGAGCAGTGGACCAATGTAAGAAACGATCCAAACGTTTCGAGATCTG  
ATCTTCATCATAACCACCGTCTCACTGAGGAAGAGTATCTAGCTTTCTGTCTCATGCTTCTT  
GCTCGGGATGGCGGCGATCTTGACTCTGTGACGGTTGCGGAGAAGCCGAGTTATAAGTGTGG  
CGTTTGTTACAAGACGTTTTCGTCTTACCAAGCTCTCGGCGGTCATAAAGCGAGCCACCGGA  
GCTTATACGGTGGTGGAGAGAATGATAAATCGACACCATCCACCGCCGTGAAATCTCACGTT  
TGTTTCGGTTTTCGGGGAAATCTTTCGCCACCGGTCAAGCTCTCGGCGGCCACAAGCGGTGCCA  
CTACGATGGTGGCGTTTCGAACTCGGAAGGTGTGGGGTCTACTAGCCACGTCAGCAGTAGTA  
GCCACCGTGGATTGACCTTAATATTATACCGGTGCAGGGATTTTCGCCGGACGACGAAAGTG  
ATGAGTCCGATGGCGACTAAGAAGCCTCGCCTGAAGTAAGTCTTTGTTGAAGACCTGGAAGT  
TTATCAAATGTAAATATCAAATTTCAATTTCAAGGAACAGTTTGTGATTCTATTACCAAT  
ACACAATACGATTCAATTCC

**SEQ ID NO 33:** Translation of gi\_30694224\_ref\_NM\_123683.2

MALEALNSPRLVEDPLRFNGVEQWTKCKKRSKRSRSDLHHNHRLTEEEYLAFCLIMLLARDGG  
DLDSVTVAEKPSYKCGVCYKTFSSYQALGGHKASHRSLYGGGENDKSTPSTAVKSHVCSVCG  
KSFATGQALGGHKRCHYDGGVSNSEGVGSTSHVSSSSHRGFDLNIIPVQGFSPDDEVMSMA  
TKKPRLK

FIGURE 3 (continued)

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SEQ ID NO 34: gi\_30698307\_ref\_NM\_126145.2 mRNA

CACACTTCACTCTTTCTTCATCTTCTTCTTAAATAGCTCGAAATCACATCTCACAGAAT  
TAAATCTTATGGCTCTCGAGACTCTCAATTCTCCAACAGCTACCACCACCGCTCGGCCTCTT  
CTCCGGTATCGTGAAGAAATGGAGCCTGAGAATCTCGAGCAATGGGCTAAAAAGAAAACGAAC  
AAAACGTCAACGTTTTGATCACGGTCATCAGAATCAAGAAACGAACAAGAACCTTCCTTCTG  
AAGAAGAGTATCTCGCTCTTTGTCTCCTCATGCTCGCTCGTGGCTCCGCCGTACAATCTCCT  
CCTCTTCTCCTCTACCGTCACGTGCGTCACCGTCCGATCACCGAGATTACAAGTGTACGGT  
CTGTGGGAAGTCCTTTTCGTCATACCAAGCCTTAGGTGGACACAAGACGAGTCACCGGAAAC  
CGACGAACACTAGTATCACTTCCGGTAACCAAGAACTGTCTAATAACAGTCACAGTAACAGC  
GGTTCGGTTGTTATTAACGTTACCGTGAACACTGGTAACGGTGTTAGTCAAAGCGGAAAGAT  
TCACACTTGCTCAATCTGTTTCAAGTCGTTTTCGCTCTGGTCAAGCCTTAGGTGGACACAAAC  
GGTGTCACTATGACGGTGGCAACAACGGTAACGGTAACGGTAAGTAGCAGCAACAGCGTAGAA  
CTCGTCTGCTGGTAGTGACGTGACGATGTTGATAATGAGAGATGGTCCGAAGAAAGTGCGAT  
CGGTGGCCACCGTGGATTTGACCTAAACTTACCGGCTGATCAAGTCTCAGTGACGACTTCTT  
AACGTTGACTGAGTTTGAGGAAAAAGTCAACTATCAAGCGAAGAAAGGGTTAGTGGACGGTG  
AAGATTAACGGTCGTTTCTTCCAGTTGCTTCGGTTTGAGCTTGACTGGGTCTGTAATGAAA  
ATGATTGGAGTGGACTTGGCATTATTATTATTATTTTAAAAAGAAATGTTAATTTGTTGTT  
GGATTTGTTTATAGATAGAGGAAACAATTGGGATACACAAATATTTTTTTTTTTTACAAAGA  
AAATAATAATGCAGAGATGGATGATTGGATCGTACACGTTATTATATAGTGGACCATTCTGT  
AATCGTGAATTATTATTATTGTTAGAAATTTAATTTTCGT

SEQ ID NO 35: Translation of gi\_30698307\_ref\_NM\_126145.2

MALETLSPTATTTARPLLRYREEMEPENLEQWAKRKRTKRQRFDHGHQONQETNKNLPSEEE  
YLALCLLMLLARGSAVQSPPLPPLPSRASPSDHRDYKCTVCGKSFSSYQALGGHKTSRHKPTN  
TSITSGNQELSNNSHNSGVSVINVTVNTGNVVSQSGKIHTCSICFKSFASGQALGGHKRCH  
YDGGNNGNGNGSSNSVELVAGSDVSDVDNERWSEESAIGGHRGFDLNLNLPADQVSVTTS

## OTHER GENES IN EVALUATION

SEQ ID NO 36: gi\_12698881\_ref\_AF\_332876.1 mRNA, 2xC2H2,

Oryza sativa

AATTCGGCACGAGCCACACAGCAACCAGCCAGCTGCCACACTAGCTTGAGGCGAGCGAGCG  
AAGCTTAGCTAGCGGATAGAACAAGTCGTCGATCTGCTTGCTGCTTTTGTGAATTGCGGTGG  
AAGCATGTGAGCGCGTCGTCCATGGAAGCGCTCCACGCCGCGGTGCTCAAGGAGGAGCAGC  
AGCAGCACGAGGTGGAGGAGGCGACGGTCGTGACGAGCAGCAGCGCCACGAGCGGGGAGGAG  
GGCGGACACCTGCCCCAGGGGTGGGCGAAGCGGAAGCGGTGCGCGCCGCCAGCGATCGGAGGA  
GGAGAACCTCGCGCTCTGCCTCCTCATGCTCGCCCGCGGCGGCCACCACCGCTCCAGGCGC  
CGCCTCCGCTCTCGGCTTCGGCGCCCCCGCCGGCAGGTGCGGAGTTCAAGTGCTCCGTCTGC  
GGCAAGTCCTTCAGCTCCTACCAGGCGCTCGGCGGCCACAAGACGAGCCACCGGGTCAAGCT  
GCCGACTCCGCCCCGAGCTCCCGTCTTGGCTCCCGCCCCCGTCCGCGCCTTGCTGCCTTCCG  
CCGAGGACCGCGAGCCAGCCACGTATCCACGCGCGCTCCTCCGACGGCATGACCAACAGA  
GTCCACAGGTGTTCCATCTGCCAGAAGGAGTTCCCGACCGGCGAGGCGCTCGGCGGGCACAA  
GAGGAAGCACTACGACGGTGGCGTAGGCGCGGCGCGCGCATCTTCAACCGAGCTCCTGG  
CCACGGTGGCGCGAGTCCGAGGTGGGAAGCTCCGGCAACGGCCAGTCCGCCACCGGGCG  
TTCGACCTCAACCTCCCGGCGGTGCCGGAGTTCGTGTGGCGGCGGTGCTCCAAGGGCAAGAA  
GATGTGGGACGAGGAGGAGGAGGTCCAGAGCCCCCTCGCCTTCAAGAAGCCCCGCTTCTCA  
CCGCGTAATTCAGCAGCTGCACGGATCCGATCCGTGAGAGTTTTGTCTAGGGAGTGAAATT  
CAGTCGAAACACACTATTCTGTTGATTCGTTTTGTGCGGCTATTGTTTAATTTGTTCTCTGCTT

FIGURE 3 (continued)

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TTGTACAGAGCAAGCGAGTGATACATAGCCATACATACAGTCATACAGATATAGGTCTAGCT  
CTTCTTGTTCTTTGTAACTGGAAGTGTACCTGTATCTTTTACACTTTGTTCTTTGACA  
GTCATATATTGTAGACCAAAAAAAAAAAAAAAAAAAAA

SEQ ID NO 37: gi\_12698882\_ref\_AAK01713.1, 2xC2H2, *Oryza sativa*  
MSSASSMEALHAAVLKEEQQHEVEEATVVTSSSATS GEEGGHLPQGWAKRKR SRQRSEEE  
NLALCLMLLARGGHRVQAPPPLSASAPPPAGAEFKCSVC GKSFSYQALGGHKTS HRVKLP  
TPPAAPVLAPAPVAALLPSAEDREPATSSTAASSDGMTNRVHRC SICQKEFPTGQALGGH KR  
KHYDGGVGAGAGASSTELLATVA AESEVGSSNGQSATRAFDLNLPAVPEFVWRPCSKGKKM  
WDEEEEVQSPLAFKKPRLTA

SEQ ID NO 38: gi\_6434215\_ref\_AL132966.1\_region 116202 ...  
116729, 2xC2H2, *Arabidopsis thaliana*

ATGAAGAGAGACCGGTCCGATTACGAAGAATCCATGAAGCATATAGACATAGTAGAAAGTCT  
AATGATGTTATCTCGAAGTTTCGTGGTCAAACAAATCGATGTAAAGCAATCTACCGGAAGCA  
AAACGAACCATAATAACCACTTCGAATGCAAAACGTGTAACCGGAAATTTGATTCTCTCAA  
GCTCTTGGAGGTCATAGAGCTAGCCACAAGAAACCTAAGCTGATCGTTGACCAAGAACAGGT  
GAAGCATCGTAACAAAGAGAATGATATGCATAAGTGTACAATTTGCGATCAAATGTTTGGGA  
CCGGTCAAGCTCTAGGCGGTCACATGAGAAAGCATAGGACGAGCATGATAACCGAGCAATCG  
ATTGTCCCTTCTGTGGTTTATTCCAGACCGTTTTTAATCGTTGCAGTAGCAGCAAGGAGAT  
CTTGGA CTTAAATCTAATCTCCATTGGAAAATGATCTTGTGTTAATCTTTGGGAAGAATTGG  
TTCCACAAATTGATTTGAAGTTTGTGAATTAG

SEQ ID NO 39: gi\_6729511\_ref\_CAB67667.1, 2xC2H2, *Arabidopsis thaliana*  
MKRDRSDYEESMKHIDIVESLMMLSRSEFVVKQIDVKQSTGSKTNHNNHFECKTCNRKFDSFQ  
ALGGHRASHKKPKLIVDQEQVKHRNKENDMHKCTICDQMFGTGQALGGHMRKHRTSMITEQS  
IVPSVVYSRPVFNRCSSESKEILDNLNLTPLNDLVLF GKNLVLPQIDLKFN

SEQ ID NO 40: ref\_CA279020, 2xC2H2, sugar cane  
CCTAACCAGCATTAGCTTTTCAAATCAACAAGCCTCGCCGTGACCGATCGATGGCCATCACC  
CAGGACGACTACGTCTCCCTCTGCCTCATGGCGCTCGCAGCCGCGGGAGGCGGAGGCCAAGC  
TGGTTTAAACAACGCAGTACGCTCTGAACACGGCTGCCTGGACAGCGACGGCGCAAGAGTCCG  
AGCTCCGCTTCCGGTGCTCCGTCTGTGGCAAGGCCTTCGCGTCGCACCAGGCACTGGGCGGG  
CACAAGGCCAGCCACCGCAAGCCGACGCTCGTACAGGCACATGCGTCGTCTCAGCCGGAGG  
CGCGGCGTCGTCTCGGTAAACAATGACCTCGGCCGTAGGCAGCAGTGGGCAGGGGAGGCACA  
GGTGACGGTGTGCCATCGGAGCTTCGCGACGGNGCAAGCGCTCGGCGGGCACAAAGAGGTGC  
CATTACTGGGACGGGCTCTCGGTGTGCTCACC GCGTCGTCTCGGCGCCATCGGGGTCCGGGTC  
GACCGTCAAGGGCTTTGATCTGAATTTGGTGCCGGTGCCGCCCCGCGATGGCCGCCAACGCTG  
CGACAAGGTGGGGAGAGGAGAAANNAAGTCANAAACCTTGGCGGTCAAGAGAAGGCGGCTTG  
CCGGTCCGTCTTGGACCCTAATTTAACGATTTAGAAAGTCCTTTTTTTAATAATTAAGAGTTC  
TTTTGAAGAAGGTTGTAAAGTTTTCGAACCTTGTCTTTTAAATGGATTGGGTGCTGGCGAA  
ATTTTAAACTGGATTAAATTTGCGCTCACTCTTTTTTTTATTTTTTACACCTTTTTTTT  
TTTTTAGAAGAAGA

FIGURE 3 (continued)

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SEQ ID NO 41: gi\_18027011\_ref\_AF254447.1, 2xC2H2, *Arabidopsis thaliana*  
TTCTTTCTCTCTCTCTCTCTCTCTTACCATTGACTGATCCTTATTCCAATTTCTTCACA  
GACTGGTTCAAGTCTAATCCTTTTACCATTACCCTAATTCCTCCACTAACCCTCTCCTCA  
TCCTCTTCTCTCTGTTACTCCTCCCTCTTCTCTTCTCTTCTTCCCTCAATCCGGAGACCTCC  
GCCGTCCACCGCCGCCACCAACTCCTCCTCTCTCTCTCTCTCCGAGAAGCCCTCCCTCTC  
CTCAGCCTCAGCCCCGCCAACAAACAACAAGACCACCATCACAACCATGACCACCTTATTCA  
AGAACCACCTTCAACCTCCATGGATGTCGACTACGATCATCACCATCAAGATGATCATCATA  
ACCTCGATGACGATGACCATGACGTCACCGTTGCTCTTCACATAGGCCTTCCAAGCCCTAGT  
GCTCAAGAGATGGCCTCTTTGCTCATGATGTCTTCTTCTTCTCTTCTCTCGAGGACCACTCA  
TCATCACGAGGACATGAATCACAAGAAAGACCTCGACCATGAGTACAGCCACGGAGCTGTCTG  
GAGGAGGAGAAGATGACGATGAAGATTGAGTCGGCGGAGACGGCGGCTGTAGAATCAGCAGA  
CTCAACAAGGGTCAATATTGGATCCCTACACCTTCTCAGATTCTCATTTGGCCCTACTCAGTT  
CTCATGTCTGTTTGTCTTCAAACCTTCAACAGATACAATAACATGCAGATGCATATGTGGG  
GACATGGATCACAATACAGAAAAGGACCTGAATCTCTAAGGGGAACACAACCAAGGAATG  
CTAAGGCTTCCGTGCTATTGCTGCGCCCCAGGCTGTCGCAACAACATTGACCATCCAAGGGC  
AAAGCCTCTCAAAGACTTCAGAACCCTTCAAACACATTACAAGAGAAAACATGGGATCAAAC  
CTTTTCATGTGTAGGAAATGTGGAAAGGCTTTTCGCGAGTCCGAGGGGACTGGAGAACACATGAG  
AAGAATTGTGGCAAACCTTTGGGTATTGCATATGTGGATCTGATTTCAAGCACAAGAGATCTCT  
CAAAGATCACATCAAGGCTTTTGGGAATGGTCATGGAGCCTACGGAATTGATGGGTTTGATG  
AAGAAGATGAGCCTGCCTCTGAGGTAGAACAATTAGACAATGATCATGAGTCAATGCAGTCT  
AAATAGCTTATATATATTACTATAAGTACTAAGTAATTCGGTATATATATTAATTATAAGAA  
ACCTAAATCTATGGACCAAGTTTTGATGGAGGTAGGGCTTTTCAAACCTAAAAGCTATATCAT  
CTAATTGATCATAGGAAAAAATGAATCAAGAGCACTTGGAAAAATTTAAATTGTATCTTTA  
GCTTCTTAGTTAAATTTATTGCAAGACAATGTAGCAGTCTAACCAATGAGGTTCCCAACGGT  
TTATTTCTATTTGTATATTATTTGTTCATTAGCTTCACCTTTCGTTAATTGAAGGACATAA  
CTTATAAATGTTTAAATTATG

SEQ ID NO 42: At3g57670, 2xC2H2, *Arabidopsis thaliana*  
MTDPYSNFFTDWFKSNPFHHYPNSSTNPSPHLPVTPPSSFFFFPQSGDLRRPPPPPTPPP  
SPPLREALPLLSLSPANKQQDHHHNDHLIQEPPSTSMVDVYDHHHQDDHHNLDDDDHDVTV  
ALHIGLPSPSAQEMASLLMSSSSSSSRTHHHEDMNHKKDLDEYSHGAVGGGEDDDDEDSV  
GGDGGCRISRLNKQYWIPTPSQILIGPTQFSCPVCFKTFNRYNNMQMHMWGHGSQYRKGP  
SLRGTOPTGMLRLPCYCCAPGCRNNIDHPRAKPLKDFRTLQTHYKRKHGKIPFMCRCGKAF  
AVRGDWRTHKNCGLWYICGSDFKHKRSLKDHIAFGNGHGAYGIDGFDEEDEFASEVEQ  
LDNDHESMQSK

SEQ ID NO 43: gi\_18676370\_ref\_AJ311810.2, 2xC2H2, *Arabidopsis thaliana*  
ATCTACACACTACTACTCACATCTCATCTCTCTCTAGCACATACCCATCAAACCATATAGAT  
ACGGTGCTTTTATTCTTGATCTTCTTCTTCTTTGTCTTCTCCTCAGAGTCATGTCTAAT  
CCAGCTTGTCGAATCTCTTCAACAATGGATGTGACCATAATAGCTTCAACTATTCCACTTC  
TCTCTCTTACATTTACAACCTCTCACGGTAGCTACTATTACTCTAATACCACAAACCTAATT  
ACATTAATCATACTCATACCACTTCCACTTCCCCTAACTCACCCCCACTAAGAGAAGCTCTT  
CCTCTTCTTAGCTTAAGCCCCATAAGGCACCAAGAACAACAAGACCAACACTATTTTCATGGA  
CACCCATCAAATTAGCTCTTCAAACCTTCTTGATGATCCTCTTGTGACTGTGGATCTTCATC  
TAGGGTTACCAAACCTACGGTGTTGGTGAGAGCATTAGGAGCAATATTGCTCCTGATGCAACC

FIGURE 3 (continued)

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ACGGACGAGCAAGATCAAGATCATGACCGAGGAGTAGAAGTCACAGTTGAGTCCCACCTTGA  
 TGATGATGATGATCATCATGGAGATCTACACAGAGGTCATCACTATTGGATTCCCTACTCCTT  
 CTCAGATTTTGATTGGTCCTACACAGTTCACTTGTCTCTTTGCTTCAAGACATTC AACAGA  
 TACAACAACATGCAGATGCACATGTGGGGACACGGCTCACAAACAGAAAGGGACCAGAATC  
 CTTAAGAGGAACCCAACCAACAGGAATGCTAAGACTACCATGTTTCTGCTGTGCACCCGGTT  
 GCAAGAACAACATTGACCACCCACGAGCCAAGCCTCTTAAGGACTTTCGAACCCCTCCAAACA  
 CATTACAAACGTAAACATGGGTCTAAACCATTGTGCTTGTCTGATGTGTGGTAAGGCCTTTGC  
 AGTGAAAGGAGATTGGAGAACGCATGAGAAGAATTGTGGAAAGCTTTGGTATTGCTCTTGTG  
 GCTCGGATTTTAAGCACAAGAGGTCGCTTAAGGACCATGTCAAGGCCTTTGGAAATGGTCAT  
 GTTCCTTGTGGGATTGATAGTTTGGAGGAGATCATGAGGACTACTATGATGCTGCTTCTGA  
 TATCGAGCAATAAGATGATAGCAACAACAATGAGTGTTAATTAGGGGTTTTGTTTATTTTTC  
 CTCTCATGCATTAGTTGATTGTATGCACGTGTTCTTTAGTTTTGTTCTTCGGATCTTTGTTT  
 TATTTTGTGTTTGAGCTGTTTTTTTTTTAATTACTAAGAAGTTAATTATCATCTAAAGATTTT  
 C

**SEQ ID NO 44:** gi\_18376498\_ref\_CAC86167.1, 2xC2H2, Arabidopsis  
 thaliana  
 MSNPACSNLFNNGCDHNSFNYSTLSYIYNHSGSYYSNTTNPYINHTHTTSTSPNSPPLR  
 EALPLLSLSPIRHQEQDQHYFMDTHQISSNFDLDDPLVTVDLHLGLPNYGVGESIRSNIAP  
 DATTEQDQDHRGVEVTVESHLDDDDHHDHLDHRGHHYWIPTPSQILIGPTQFTCPCLCFKT  
 FNRYNNMQMHMWHGHSQYRKGPESLRGTQPTGMLRLPCFCCAPGCKNNIDHPRAKPLKDFRT  
 LQTHYKRKHGSKPFACRMCGKAFVKGDWRTHEKNCGLWYCSCGSDFKHKRSLKDHVKAFF  
 NGHVPCGIDSFGGDHEDYDAASDIEQ

**SEQ ID NO 45:** gi\_7798991\_ref\_AL355775.1\_region 7957 ... 8451,  
 2xC2H2, Arabidopsis thaliana  
 ATGGTTGCGAGAAGTGAGGAAGTTGAGATAGTGGAAGATACGGCGGCGAAATGTTTGATGTT  
 GTTATCAAGAGTTGGAGAATGCGGCGGAGGAGAGAAACGAGTTTCCGATGCAAGACTT  
 GTCTTAAAGAGTTTTCGTCGTTTCAAGCTTTGGGAGGTCATCGTGCAAGCCACAAGAACTC  
 ATTAACAGTAGCGATCCATCACTTCTTGGATCCTTGTCTAACAAGAAAACATAAACGGCGAC  
 GTCTCATCCTTGTCCGATATGTGGCGTGGAGTTTCCGATGGGGCAAGCTCTTGGTGGTCACA  
 TGAGGAGACATAGGAGTGAGAAAGCCTCACCAGGCACGTTGGTTACACGTTCTTTTTTACCG  
 GAGACGACGACGGTGACGACTTTGAAAAAATCGAGTAGTGGAAGAGAGTGGCTTGTTTGA  
 CTTAGATTCGATGGAGAGTTTAGTCAATTGGAAGTTGGAGTTGGGAAGAAGATTCTTGA

**SEQ ID NO 46:** gi\_7798996\_ref\_CAB90935.1, 2xC2H2, Arabidopsis  
 thaliana  
 MVARSEEVEIVEDTAAKCLMLLSRVGECGGGGEKRVFRCKTCLKEFSSFQALGGHRASHKKL  
 INSSDPSLLGSLSNKKTKTATSHPCPICGVEFPMGQALGGHMRHRSEKASPGTLVTRSFLP  
 ETTTVTTLKKSSSGKRVACLDLDSMESLVNWKLELGRITIS

**SEQ ID NO 47:** gi\_9755794\_ref\_AL391143.1\_region 31730 ...  
 32938, 2xC2H2, Arabidopsis thaliana  
 ATGGAAGACGAACATCAAGATCTCCATAAACCCATTAATGGAGCTTTGCGAGACCTCAAGAT  
 TACTCGGTACAGAAAGAAACAGAAAAGTCTACGAACCAACAGCAAGATGTTACTTGTTACT  
 ATGGTCTAAGGGAAAACCTCGAAGAAGAAAACCCAGGAATCTCCGGAACCAATGAAGAAGATT  
 TTGTTTCGATGCGAAGAATGTGGAAAAGGGTTTCGGTACGAGAAATATTTTAAGAATCATCG

FIGURE 3 (continued)

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CTCGATGATGCATTTATCGCCGAACGAGAAGGTTTGTGAAGAATCCTTGATGACTCTGTCTC  
GTAGCCTTGGGTTTGTGAAGAAGAAGAAAAGATCAAGACTTGGTAGGTCTGGGAAGACTTTA  
TTTACTACGTTTCTTGAACCGAGTTCTATTTTTGATGCGACTGATGAAGAATTAGAAGTGGC  
GGATTGTTTGTATTCTATTGTCTAAGAGTGTCTCCAAGGTTGTAGACGAATTGAAAAGTCTTT  
CTGAGGCAGTACGTGTTACTCCTGAAACACCTGAAAGTAGCTATGATTGGGTTGTTTGCTC  
AACAAGAAACCGAGAAAAGGTGGTGAATTGGAATCTGGGGTTTTAAGTAATGAGCAAAGACT  
TATGGAAGAAGGTTTTAGTAGTTATGGAACATCGAAAGAACCAGCTAGCTTCTTGAGAGACG  
AAAACAGATTGGATCAGCAGAAACGGAGAAAAGATGGTGAATTTGAATCCGGACTTTTGAGT  
AATGAGCAAAGACTGCTAGAAGAAGAGATTACTACTCCTGTGACATTCAAAGGTCCAGCGAG  
TTCCTTGAGACACAAGTGTGCTTTGGATCGAAATGGAGGTGAATTTGGTCTGAGTTTTTGA  
GTAATGAGCAAACACTGATGGAAGAAACATGGAAAGAACCAGTGAGTTTCTTAGAAGATAAG  
CATGAATTTGATCAGCGGAAAATGCGAGAAGCTGGCGACTTTGAATCTAGGTTTTACAGAAT  
TGAGCTTGGAGTAGGAGCTATGGAGTGTACTTCTTCAGATACTGATATGCTCACGCAATCTG  
ATAAGAAGAACGTTGAGCATCGATGCAGGTTGTGCAACAAGATATTCTCGTCTTATCAAGCT  
CTAGGGGGTTCATCAGACGTTTTCATCGGATGAGCAAATGTAAGAACAAGAAGAATGGCATAGA  
GGAATCAGTTGAACCCAGGATGACTCTGTGA

SEQ ID NO 48: gi\_9755803\_ref\_CAC01747.1, 2xC2H2, Arabidopsis  
thaliana

MEDEHQDLHKPINGALRDLKITRSQKETEKSTNQQQDVTCYYGLRENSKKKTQESPEPMKKI  
LFRCEECGKGFYRYEKFKNHRSMMHLSPNKEVCEESLMTLSRSLGFVKKKKRSRLGRSGKTL  
FTTFLEPSSIFDATDEELEVADCLILLSKSPKVVDLKSLSLSEAVRVTPETPESYDLGCLL  
NKKPRKGGELESGVLSNEQRLMEEGFSSYGTSEKFPASFLRDENRLDQQRKRDGEFESGLLS  
NEQRLLEEEITTPVTFKGPASSLRHKCALDRNGGEFGPEFLSNEQTLMEETWKEPVSFLEDK  
HEFDQRKMREAGDFESRFYRIELGVGAMECTSSDTDMLTQSDKKNVEHRCRLCNKIFSSYQA  
LGGHQTFHRMSKCKNKNKNGIEESVEPRMTL

SEQ ID NO 49: gi\_1418338\_ref\_X98678.1, 2xC2H2, Arabidopsis  
thaliana

CTTGTTAGTTCACTCCACATAATAAACACCAAAGATTTCAATTCTCTTCTCCATAATTTTCGAA  
GTTTCTTGAATTGGGTTTGTCTTCTGATTTGTTTCTTGAATTGGGTTTGGTCTTCTTTTCT  
TACTATATTTGGATATGATGATGGGTCAAGATGAGGTTGGGAGTGATCAGACGCAAATCATA  
AAAGGGAAACGTACGAAGCGACAAAGATCGTCTTCGACGTTTGTGGTGACGGCGGCGACAAC  
AGTGACTTCAACAAGTTCATCGGCCGGTGGAAGTGGAGGAGAAAGAGCTGTTTCAGATGAAT  
ACAACCTCGGCGGTTTCGTCTCCGGTGACTACTGATTGTACGCAAGAAGAAGACATGGCG  
ATTTGTCTCATCATGTTAGCTCGTGGGACAGTTCTTCCATCGCCGGATCTCAAGAACTCGAG  
AAAAATTCATCAGAAGATTTTCGTGGGAGAAATTCAGTTTCTATGTGTACGAGTGTAACCGT  
GTAACCGGACGTTTTTCGTGTTCCAAGCACTTGGTGGACACAGAGCGAGCCACAAGAAGCCG  
AGGACGTCGACTGAGGAAAAGACTAGACTACCCCTGACGCAACCCAAGTCTAGTGCATCAGA  
AGAAGGGCAAAACAGTCATTTCAAAGTTTCCGGCTCAGCCCTAGCTTCACAGGCAAGTAACA  
TCATCAACAAGGCAACAAAGTACACGAGTGTTCATCTGCGGTTCTGAGTTCACTTCCGGG  
CAAGCTCTCGGTGGTCACATGAGGCGGCACAGGACAGCCGTAACCACGATTAGCCCCGTTGC  
AGCCACCGCAGAAGTAAGCAGAAACAGTACAGAGGAAGAGATTGAGATCAATATAGGCCGTT  
CGATGGAACACAGAGGAAATATCTACCGTTGGATCTTAATCTACCAGCACCAGGAGATGAT  
CTAAGAGAGTCCAAGTTTCAAGGGATAGTATTCTCAGCAACACCAGCGTTAATAGATTGTCA  
TTACTAGTTGTTTTTTTTTACTACATAATATGATGAAATATTTGTGAATTCTTCTTACTTACT  
ACTATATTGTTGATCAAAAAAAAAAAAAAAAAA

FIGURE 3 (continued)

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SEQ ID NO 50: gi\_1418339\_ref\_CAA67236.1, 2xC2H2, Arabidopsis  
thaliana  
MGQDEVGSDQTQIIKGKRTKRQRSSSTFVVTAATTVTSTSSSAGGSGGERAVSDEYN SAVSS  
PVTTDCTQEEEDMAICLIMLARGTVLPSPDLKNSRKIHQKISSENSSFYVYECKTCNRTFSS  
FQALGGHRASHKKPRTSTEEKTRLPLTPKSSASEEGQNSHFKVSGSALASQASNIINKANK  
VHECSICGSEFTSGQALGGHMRHRHTAVTTISPVAATAEVSRNSTEEEEIEINIGRSMEQQRK  
YLPLDLNLPAPGDDLRESKFQGIVFSATPALIDCHY

FIGURE 3 (continued)

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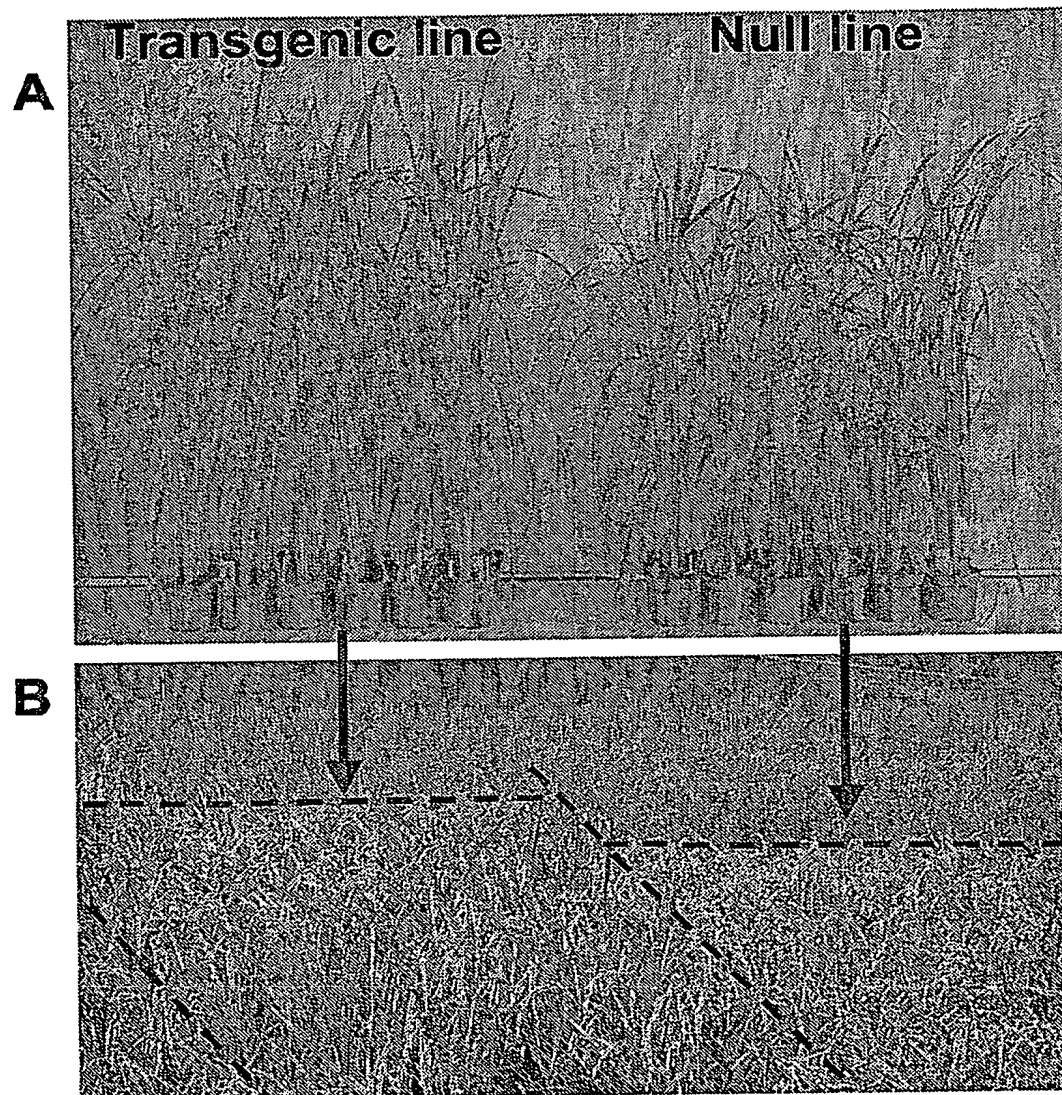


FIGURE 4



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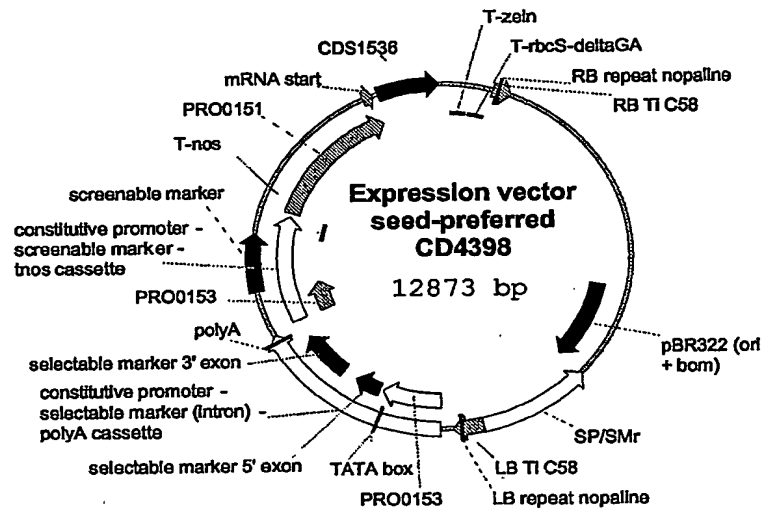


FIGURE 5